

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/596,784

TECH CENTER 1600

DATE: 11/01/2000  
TIME: 03:02:30

INPUT SET: S36052.raw

This Raw Listing contains the General  
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

(1) General Information:

(i) APPLICANT: Bogdanove, Adam J.  
Kim, Jihyun Francis  
Wei, Zhong-Min  
Beer, Steven V.

(ii) TITLE OF INVENTION: HYPERSENSITIVE RESPONSE ELICITOR FROM  
ERWINIA AMYLOVORA, ITS USE, AND ENCODING GENE

(iii) NUMBER OF SEQUENCES: 5

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP  
(B) STREET: P.O. Box 1051, Clinton Square  
(C) CITY: Rochester  
(D) STATE: New York  
(E) COUNTRY: U.S.A.  
(F) ZIP: 14603

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk  
(B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 09/596,784  
(B) FILING DATE:  
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 09/120,663  
(B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Goldman, Michael L.  
(B) REGISTRATION NUMBER: 30,727  
(C) REFERENCE/DOCKET NUMBER: 19603/1661

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (716) 263-1304  
(B) TELEFAX: (716) 263-1600

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/596,784DATE: 11/01/2000  
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47  
48 (2) INFORMATION FOR SEQ ID NO:1:  
49  
50 (i) SEQUENCE CHARACTERISTICS:  
51 (A) LENGTH: 5517 base pairs  
52 (B) TYPE: nucleic acid  
53 (C) STRANDEDNESS: single  
54 (D) TOPOLOGY: linear  
55  
56 (ii) MOLECULE TYPE: DNA (genomic)  
57  
58  
59  
60  
61 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:  
62  
63 ATGGAATTAA AATCACTGGG AACTGAACAC AAGGCGGCAG TACACACAGC GGCGCACAAAC 60  
64  
65 CCTGTGGGGC ATGGTGTTC CTTACAGCAG GGCAGCAGCA GCAGCAGCCC GCAAAATGCC 120  
66  
67 GCTGCATCAT TGGCGGCAGA AGGCAAAAAT CGTGGGAAAA TGCCGAGAAT TCACCAGCCA 180  
68  
69 TCTACTGCGG CTGATGGTAT CAGCGCTGCT CACCAGCAAA AGAAATCCTT CAGTCTCAGG 240  
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71 GGCTGTTTGG GGACGAAAAA ATTTTCCAGA TCGGCACCGC AGGGCCAGCC AGGTACCACC 300  
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73 CACAGCAAAG GGGCAACATT GCGCGATCTG CTGGCGCGGG ACGACGGCGA AACGCAGCAT 360  
74  
75 GAGGCGGCCG CGCCAGATGC GGCGCGTTTG ACCCGTTCGG GCGGCGTCAA ACGCCGCAAT 420  
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77 ATGGACGACA TGGCCGGGCG GCCAATGGTG AAAGGTGGCA GCGGCGAAGA TAAGGTACCA 480  
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81 AAAATGGCTC ACCCGGCTTC AGCCAACGCC GGCGATCGCC TGCAGCATTC ACCGCCGCAC 600  
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83 ATCCCGGGTA GCCACCACGA AATCAAGGAA GAACCGGTTG GCTCCACCAG CAAGGCAACA 660  
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85 ACGGCCCACG CAGACAGAGT GGAAATCGCT CAGGAAGATG ACGACAGCGA ATTCCAGCAA 720  
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87 CTGCATCAAC AGCGGCTGGC GCGCGAACGG GAAAATCCAC CGCAGCCGCC CAAACTCGGC 780  
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89 GTTGCCACAC CGATTAGCGC CAGGTTTCAG CCCAACTGA CTGCGGTTGC GGAAAGCGTC 840  
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91 CTTGAGGGGA CAGATACCAC GCAGTCACCC CTTAAGCCGC AATCAATGCT GAAAGGAAGT 900  
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93 GGAGCCGGGG TAACGCCGCT GGCGGTAACG CTGGATAAAG GCAAGTTGCA GCTGGCACCG 960  
94  
95 GATAATCCAC CCGCGCTCAA TACGTTGTTG AAGCAGACAT TGGGTAAAGA CACCCAGCAC 1020  
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97 TATCTGGCGC ACCATGCCAG CAGCGACGGT AGCCAGCATC TGCTGCTGGA CAACAAAGGC 1080  
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99 CACCTGTTTG ATATCAAAAG CACCGCCACC AGCTATAGCG TGCTGCACAA CAGCCACCCC 1140

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PATENT APPLICATION US/09/596,784DATE: 11/01/2000  
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155	CAAACCGTCT TTAACCGACT AATGCAGGGG GTGAAAGGCA AGGTGATCCC AGGCAGCGGG	2820
156		
157	TTGACGGTTA AGCTCTCGGC TCAGACGGGG GGAATGACCG GCGCCGAAGG GCGCAAGGTC	2880
158		
159	AGCAGTAAAT TTTCCGAAAG GATCCGCGCC TATGCGTTCA ACCCAACAAT GTCCACGCCG	2940
160		
161	CGACCGATTA AAAATGCTGC TTATGCCACA CAGCACGGCT GGCAGGGGCG TGAGGGGTTG	3000
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163	AAGCCGTTGT ACGAGATGCA GGGAGCGCTG ATTAAACAAC TGGATGCGCA TAACGTTCTG	3060
164		
165	CATAACGCGC CACAGCCAGA TTTGCAGAGC AAAGTGGAAA CTCTGGATTT AGGCGAACAT	3120
166		
167	GGCGCAGAAT TGCTTAACGA CATGAAGCGC TTCCGCGACG AACTGGAGCA GAGTGCAACC	3180
168		
169	CGTTCGGTGA CCGTTTTAGG TCAACATCAG GGAGTGCTAA AAAGCAACGG TGAAATCAAT	3240
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171	AGCGAATTTA AGCCATCGCC CGGCAAGGCG TTGGTCCAGA GCTTTAACGT CAATCGCTCT	3300
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173	GGTCAGGATC TAAGCAAGTC ACTGCAACAG GCAGTACATG CCACGCCGCC ATCCGCAGAG	3360
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175	AGTAAACTGC AATCCATGCT GGGGCACTTT GTCAGTGCCG GGGTGGATAT GAGTCATCAG	3420
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177	AAGGGCGAGA TCCCCTGGG CCGCCAGCGC GATCCGAATG ATAAAACCGC ACTGACCAAA	3480
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179	TCGCGTTTAA TTTTAGATAC CGTGACCATC GGTGAAGTGC ATGAAGTGGC CGATAAGGCG	3540
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181	AAAGTGGTAT CTGACCATAA ACCCGATGCC GATCAGATAA AACAGCTGCG CCAGCAGTTC	3600
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183	GATACGCTGC GTGAAAAGCG GTATGAGAGC AATCCGGTGA AGCATTACAC CGATATGGGC	3660
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193	AAGAAGGTGC CAGTTCCGGT GATCCCCGGA GCCGGCATCA CGCTGGATCG CGCCTATAAC	3960
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195	CTGAGCTTCA GTCGTACCAG CGGCGGATTG AACGTCAGTT TTGGCCGCGA CGGCGGGGTG	4020
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199	ACCAGTGCAG GTAACGCCAG TGAAGTGTG AGCGCAAAAC ATAAATCAG CCCGACTTG	4140
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201	CGTATCGGCG CTGCTGTGAG TGGCACCCTG CAAGGAACGC TACAAAACAG CCTGAAGTTT	4200
202		
203	AAGCTGACAG AGGATGAGCT GCCTGGCTTT ATCCATGGCT TGACGCATGG CACGTTGACC	4260
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205	CCGGCAGAAC TGTTCAGAAA GGGGATCGAA CATCAGATGA AGCAGGGCAG CAAACTGACG	4320

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207	GGCAGTAAAC	CAAATGGTGT	CACTGCCCCG	GTTTCTGCCG	GGCTAAGTGC	ATCGGCAAAC	4440
208	CTGGCCGCCG	GCTCGCGTGA	ACGCAGCACC	ACCTCTGGCC	AGTTTGGCAG	CACGACTTCG	4500
209	GCCAGCAATA	ACCGCCCAAC	CTTCCTCAAC	GGGGTCGGCG	CGGGTGCTAA	CCTGACGGCT	4560
210	GCTTTAGGGG	TTGCCCATTG	ATCTACGCAT	GAAGGGAAAC	CGGTGCGGAT	CTTCCCGGCA	4620
211	TTTACCTCGA	CCAATGTTTC	GGCAGCGCTG	GCGCTGGATA	ACCGTACCTC	ACAGAGTATC	4680
212	AGCCTGGAAT	TGAAGCGCGC	GGAGCCGGTG	ACCAGCAACG	ATATCAGCGA	GTTGACCTCC	4740
213	ACGCTGGGAA	AACACTTTAA	GGATAGCGCC	ACAACGAAGA	TGCTTGCCGC	TCTCAAAGAG	4800
214	TTAGATGACG	CTAAGCCCGC	TGAACAACTG	CATATTTTAC	AGCAGCATTT	CAGTGCAAAA	4860
215	GATGTCGTCG	GTGATGAACG	CTACGAGGCG	GTGCGCAACC	TGAAAAAACT	GGTGATACGT	4920
216	CAACAGGCTG	CGGACAGCCA	CAGCATGGAA	TTAGGATCTG	CCAGTCACAG	CACGACCTAC	4980
217	AATAATCTGT	CGAGAATAAA	TAATGACGGC	ATTGTGCGAG	TGCTACACAA	ACATTTTCGAT	5040
218	GCGGCATTAC	CAGCAAGCAG	TGCCAAACGT	CTTGGTGAAA	TGATGAATAA	CGATCCGGCA	5100
219	CTGAAAGATA	TTATTAAGCA	GCTGCAAAGT	ACGCCGTTCA	GCAGCGCCAG	CGTGTCGATG	5160
220	GAGCTGAAAG	ATGGTCTGCG	TGAGCAGACG	GAAAAAGCAA	TACTGGACGG	TAAGGTCGGT	5220
221	CGTGAAGAAG	TGGGAGTACT	TTTCCAGGAT	CGTAACAACT	TGCGTGTTAA	ATCGGTCAGC	5280
222	GTCAGTCAGT	CCGTCAGCAA	AAGCGAAGGC	TTCAATACCC	CAGCGCTGTT	ACTGGGGACG	5340
223	AGCAACAGCG	CTGCTATGAG	CATGGAGCGC	AACATCGGAA	CCATTAATTT	TAAATACGGC	5400
224	CAGGATCAGA	ACACCCACG	GCGATTTACC	CTGGAGGGTG	GAATAGCTCA	GGCTAATCCG	5460
225	CAGGTCGCAT	CTGCGCTTAC	TGATTTGAAG	AAGGAAGGGC	TGGAAATGAA	GAGCTAA	5517

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 1838 amino acids

(B) TYPE: amino acid

(C) STRANDEDNESS:

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

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**SEQUENCE VERIFICATION REPORT**  
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**SEQUENCE MISSING ITEM REPORT**  
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**SEQUENCE CORRECTION REPORT**  
**PATENT APPLICATION US/09/596,784**

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